

# SEQUENCE LISTING

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<120> Novel method of selecting immunosuppressant having little thrombocytopenic effect

<130> 264163US0PCT

<140> 10/519,678  
<141> 2005-01-07

<150> PCT/JP03/08621  
<151> 2003-07-07

<150> JP 2002-203901  
<151> 2002-07-12

<160> 19

<170> PatentIn version 3.1

<210> 1  
<211> 32  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: PCR primer-1

<400> 1  
tcgctagcct gagtatttaa caatcgacc ct 32

<210> 2  
<211> 30  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: PCR primer- 2

<400> 2  
cgaagcttgt ggcaggagtt gaggttactg 30

<210> 3  
<211> 30  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: PCR primer-3

<400> 3  
 cgctagctgc tcttgtccac cacaatatgc 30

<210> 4  
 <211> 28  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Description of Artificial Sequence: PCR primer-4

<400> 4  
 atagatctat ccctggctcc cacctcag 28

<210> 5  
 <211> 28  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Description of Artificial Sequence: PCR primer-5

<400> 5  
 ataagctttg gtggttgcg agggttcg 28

<210> 6  
 <211> 28  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Description of Artificial Sequence: PCR primer-6

<400> 6  
 atggtaccac cccagaagat gccaggag 28

<210> 7  
 <211> 28  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Description of Artificial Sequence: PCR primer- 7

<400> 7  
 atgctagcgc cctctgagcc tcagtttc 28

<210> 8  
 <211> 731  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> misc\_feature  
 <222> (1)..(731)

<223> Human interleukin-2 (IL-2) gene 5'-flanking region

<220>

<221> TATA\_signal

<222> (652)..(657)

<223> Corresponding to the TATA sequence

<220>

<221> misc\_feature

<222> (676)..(657)

<223> The putative transcription initiation site of the IL-2 gene promoter

<220>

<221> misc\_feature

<222> (1)..(731)

<223> Corresponding to the sequence (689-1416) in the GenBank database (Accession: X00695), except for several variations.

<220>

<221> variation

<222> (38)..(731)

<223> 1 bp insertion

<220>

<221> variation

<222> (196)..(731)

<223> T to A exchange

<220>

<221> variation

<222> (346)..(731)

<223> T to G exchange

<220>

<221> variation

<222> (577)..(731)

<223> T to A exchange

<220>

<221> variation

<222> (688)..(689)

<223> 2 bp insertion (CT)

<400> 8

ctgagtattt aacaatcgca ccctttaaaa aatgtacaat agacattaag agacttaaac 60

agatatataa tcatttttaa ttaaaatagc gttaaacagt acctcaagct caataagcat 120

tttaagtatt ctaatcttag tatttctcta gctgacatgt aagaagcaat ctatcttatt 180

gtatgcaatt agctcattgt gtggataaaa aggtaaaacc attctgaaac aggaaaccaa 240

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tacacttcoct gtttaaatcaa caaatctaaa catttattct tttcatctgt ttactcttgc      300
tcttgtccac cacaatatgc tattcacatg ttcagtgtag ttttaggaca aagaaaattt      360
tctgagttac ttttgtatcc ccaccccctt aaagaaagga ggaaaaactg tttcatacag      420
aaggcggttaa ttgcatgaat tagagctatc acctaagtgt gggctaattgt aacaaagagg      480
gatttcacct acatccattc agtcagtcct tgggggttta aagaaattcc aaagagtcac      540
cagaagagga aaaatgaagg taatgttttt tcagacaggt aaagtctttg aaaatatgtg      600
taatatgtaa aacattttga ccccccata atatttttcc agaattaaca gtataaattg      660
catctcttgt tcaagagttc cctatcactc tctttaatca ctactcacag taacctcaac      720
tcctgccaca a                                                                731

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<210> 9
<211> 819
<212> DNA
<213> Homo sapiens

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<220>
<221> misc_feature
<222> (1)..(819)
<223> Human GATA- 1 gene promoter region

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<220>
<221> misc_feature
<222> (1)..(819)
<223> Corresponding to the sequence (5342-6160) reported by K. Blechsc
      hmidt et al. / GenBank (Accession: AF196971)

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<220>
<221> misc_feature
<222> (790)..(819)
<223> The putative transcription initiation site of the GATA-1 gene pro
      moter

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<400> 9
atccctggct cccacctcag tttcccgctt ccaaggcagc atggcgggca agaagttgag      60
gccactgtcc ctgggtgttc ctacccccac accctcaccc caagacagcc tgttactgag      120
gcgccaacag ccacggtcgc ctacatctga taagacttat ctgctgcccc agggcaggcc      180
ggagctggcg taagccccag tggggcgcta agtgagtgtg ccctgcctc ccgccagcac      240
tggcctggcc tgcaggctta gcctgggtca tcaaggtatc ccacaggctc tagttcaaat      300
ccagcagaac ctctctgagc ctactcttcc tcacctgcaa aatgggtaca gccacatccc      360
ttctctccct gcagccagga agacgcacat acacaggagt ctagcccaca ccggccccgc      420
acaaattaag ggctttactc tctgaaaagc ccagtgaagt catgaaacca tatctgotat      480

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tttcatttat cttggtttca gcctatTTTtg cttgtctgga cactacagtc cacgggagcc 540
taggtcgagc gaggtccaag aatccccagg gtgggcaggg aggggtggaag agggcctcca 600
gtgccaaga ggtgccccac aagcatggga cccgccccct cccctggact gccccacca 660
ctggggcacc agccactccc tggggaggag ggaggaggga gaaggaggag agggaggag 720
ggaggaaggg agcctcaaag gccaaaggcca gccaggacac cccctgggat cacactgagc 780
ttgccacatc cccaaggcgg ccgaaccctc cgcaaccac 819

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<210> 10
<211> 637
<212> DNA
<213> Homo sapiens

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<220>
<221> misc_feature
<222> (1)..(637)
<223> Human GATA- 1 gene enhancer region

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<220>
<221> misc_feature
<222> (1)..(637)
<223> Corresponding to the sequence (2362-2998) reported by K. Blechs
      chmidt et al. / GenBank (Accession: AF196971)

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<400> 10
acccagaag atgccaggag ggagtgagcc agtcaggga ggcttccgag aagagaggac 60
attgaagaag agtctcaaac ttaggcctga cggagaagac gcgcggccag gacacccac 120
ccccgccctc gtctcccca aagcctgatc tggccccact gattccctta tctgccact 180
cccagctgcc tccttgctgg ctgaactgtc gccgcagact tctgagctg cgcgccctcc 240
acgggggatgg gggagggaat ggggtgaggc ctggcctcac agcctcgggg ttccagctc 300
ttgctggagg cagggtcttg gggcgcccta ctctcacc ttggcttctc ttctgagcg 360
ctctgtgctc tccagaaatg aagaaatggg gtgagtccag cggccaaacc cttgtcttag 420
ctcttagaca tgcctcgagc ctgccattcc ctgtgaggac agatttcct atgttgcgac 480
cgctgcttct aataataata atgatgatga taattcccat ttacagagca caccatttat 540
ggtgtgccag caggccctgt gctgagtggg tcctaccac gtggggggct aggactttac 600
ccgttttcca gatgaagaaa ctgaggctca gagggcg 637

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<210> 11
<211> 434
<212> DNA
<213> Homo sapiens

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<220>

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<221> misc\_feature  
 <222> (1)..(434)  
 <223> Human interleukin-2 (IL-2) gene 5'-flanking region

<220>  
 <221> TATA\_signal  
 <222> (355)..(360)  
 <223> Corresponding to the TATA sequence

<220>  
 <221> misc\_feature  
 <222> (379)..(360)  
 <223> The putative transcription initiation site of the IL-2 gene promoter

<220>  
 <221> misc\_feature  
 <222> (1)..(434)  
 <223> Corresponding to the sequence (+985 to +1416 ) in the GenBank database (Accession: X00695), except for several variations.

<220>  
 <221> variation  
 <222> (49)..(434)  
 <223> T to G exchange

<220>  
 <221> variation  
 <222> (280)..(434)  
 <223> T to A exchange

<220>  
 <221> variation  
 <222> (391)..(392)  
 <223> 2 bp insertion (CT)

<400> 11  
 tgctcttgtc caccacaata tgctattcac atgttcagtg tagtttttagg acaaagaaaa 60  
 ttttctgagt tactttttgta tccccacccc cttaaagaaa ggaggaaaaa ctgtttcata 120  
 cagaaggcgt taattgcatg aattagagct atcacctaag tgtgggctaa tgtaacaaag 180  
 agggatttca cctacatcca ttcagtcagt ctttgggggt ttaaagaaat tccaaagagt 240  
 catcagaaga ggaaaaatga aggtaatggt ttttcagaca ggtaaagtct ttgaaaatat 300  
 gtgtaatatg taaaacattt tgacaccccc ataatatattt tccagaatta acagtataaa 360  
 ttgcatctct tgttcaagag ttccctatca ctctctttaa tcactactca cagtaacctc 420  
 aactcctgcc acaa 434

<210> 12

<211> 59  
 <212> DNA  
 <213> Artificial Sequence  
  
 <220>  
 <223> CEBP-11, synthetic DNA  
  
 <400> 12  
 cgcgttgagc aagacttgag caagtacttg agcaagcgtt gagcaaggct tgagcaagc 59  
  
 <210> 13  
 <211> 59  
 <212> DNA  
 <213> Artificial Sequence  
  
 <220>  
 <223> CEBP-12, synthetic DNA  
  
 <400> 13  
 tcgagcttgc tcaagccttg ctcaacgctt gctcaagtac ttgctcaagt cttgctcaa 59  
  
 <210> 14  
 <211> 51  
 <212> DNA  
 <213> Artificial Sequence  
  
 <220>  
 <223> HSE-11, synthetic DNA  
  
 <400> 14  
 cgcgtctaga atgttctaga tctagaacat tctagctaga atgttctaga c 51  
  
 <210> 15  
 <211> 51  
 <212> DNA  
 <213> Artificial Sequence  
  
 <220>  
 <223> HSE-12, synthetic DNA  
  
 <400> 15  
 tcgagtctag aacattctag ctagaatggt ctagatctag aacattctag a 51  
  
 <210> 16  
 <211> 651  
 <212> DNA  
 <213> Artificial Sequence  
  
 <220>  
 <223> GATA-1 gene HSI region (mutant)  
  
 <220>  
 <221> mutation  
 <222> (178)..(178)  
 <223> in vitro mutation (from "a" to "g")

<220>  
 <221> mutation  
 <222> (519)..(519)  
 <223> in vitro mutation (from "t" to "g")

<400> 16  
 ctacgcgtac cccagaagat gccaggaggg agtgagccag tcagggaagg cttccgagaa 60  
 gagaggacat tgaagaagag tctcaaaactt aggcctgacg gagaagacgc gcggccagga 120  
 caccaccacc cgcgcctcgt ctcccccaaa gcctgatctg gcccactga ttcccttgtc 180  
 tgcccactcc cagctgcctc cttgctggct gaactgtcgc cgcagacttc tgagcctgcg 240  
 ccccctccac ggggatgggg gagggaatgg ggtgaggcct ggccacacag cctcgggggt 300  
 tccagctctt gctggaggca gggctctggg gcgcctact cctcaccctt ggcttctctt 360  
 cctgagcgcct ctgtgctctc cagaaatgaa gaaatggggg gagtccagcg gccaaaccct 420  
 tgtottagct cttagacatg cctcgagcct gccattccct gtgaggacag atttccttat 480  
 gttgcgaccg ctgcttctaa taataataat gatgatgaga attcccattt acagagcaca 540  
 ccatttatgg tgtgccagca ggccctgtgc tgagtgggtc ctaccacgt ggggggctag 600  
 gactttaccc gttttccaga tgaagaaact gaggctcaga gggcagatct g 651

<210> 17  
 <211> 838  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> GATA-1 gene IE promoter (mutant)

<220>  
 <221> mutation  
 <222> (160)..(160)  
 <223> in vitro mutation (from "t" to "g")

<220>  
 <221> mutation  
 <222> (168)..(168)  
 <223> in vitro mutation (from "a" to "g")

<220>  
 <221> mutation  
 <222> (287)..(287)  
 <223> in vitro mutation (from "a" to "g")

<220>  
 <221> mutation  
 <222> (482)..(482)  
 <223> in vitro mutation (from "t" to "g")

<220>  
 <221> mutation  
 <222> (498)..(498)  
 <223> in vitro mutation (from "a" to "g")

<220>  
 <221> mutation  
 <222> (798)..(798)  
 <223> in vitro mutation (from "t" to "g")

<400> 17  
 atagatctga tccctggctc ccacctcagt ttcccgcctc caaggcagca tggcgggcaa 60  
 gaagttgagg ccaactgtccc tgggtgttcc tccccccaca ccctcacccc aagacagcct 120  
 gttactgcgg cgccaacagc caggtgcgc tacatctgag aagacttgct tgctgcccc 180  
 gggcaggccg gagctggcgt aagccccagt ggggcgctaa gtgagtgtgc ccctgcctcc 240  
 cgccagcact ggccctggcct gcaggcttag cctgggtcat caaggtgtcc cacaggctct 300  
 agttcaaadc cagcagaacc tctctgagcc tcaactcttct cacctgcaaa atgggtacag 360  
 ccacatccct tctctccctg cagccaggaa gacgcacata cacaggagtc tagcccacac 420  
 cgcccccgca caaattaagg gctttactct ctgaaaagcc cagtgaagtc atgaaacat 480  
 agctgctatt ttcatttgct ttggtttcag cctattttgc ttgtctggac actacagtcc 540  
 acgggagcct aggtcgagcg aggtccaaga atccccaggg tgggcaggga ggggtggaaga 600  
 gggcctccag tgcccaagag gtgccccaca agcatgggac ccgccccctc ccctggactg 660  
 cccacccac tggggcacca gccactccct ggggaggagg gaggaggag aaggaggga 720  
 gggaggagg gaggaaggga gcctcaaagg ccaaggccag ccaggacacc ccctgggatc 780  
 aactgagct tgccacagcc ccaaggcggc cgaaccctcc gcaaccacca aagcttat 838

<210> 18  
 <211> 65  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> STAT3 forward sequence, synthetic DNA

<400> 18  
 tgcttccga acgttgcttc ccgaacgttg cttcccgaac gttgcttccc gaacgtagat 60  
 ctggg 65

<210> 19  
 <211> 65  
 <212> DNA  
 <213> Artificial Sequence

<220>

<223> STAT3 reverse sequence, synthetic DNA

<400> 19

cccagatcta cgttcgggaa gcaacgttcg ggaagcaacg ttcgggaagc aacgttcggg  
aagca

60

65